

Table 1

Fragment	Size	PRIMERS (5' TO 3')	LOCATION
<b>Inhibin Alpha</b>			
INH $\alpha$	601bp	For GCTGCTGCGCTGTCCCTCTGTA (SEQ ID NO:10) Rev TATTTCCCAACTCTGCCTTTCTC (SEQ ID NO:11)	732... 754 1332... 1309
INH $\alpha$ 1	243bp	For GGCCCACTCGGACCAGAC (SEQ ID NO:12) Rev AGCCACACAACCACCATGACAGTAG (SEQ ID NO:13)	792...811 1034...1011
INH $\alpha$ 2	139bp	For GCTGGGCTGGGAACGGTGGAT (SEQ ID NO:14) Rev GGAGTAGGGCTGGGCTGGGGTAGG (SEQ ID NO:15)	963...983 1101...1078
INH $\alpha$ 3	254bp	For CTACCCAGCCAGCCCTACTCCT (SEQ ID NO:16) Rev TATTTCCCAACTCTGCCTTTCTC (SEQ ID NO:17)	1079...1102 1332...1309
<b>Inhibin Beta A</b>			
INH $\beta$ A	529bp	For CTGGGCAAGAAGAAGAAGAAAGAA (SEQ ID NO:18) Rev CCTGGGCTGGGCAACTC (SEQ ID NO:19)	1005...1028 1533...1517
INH $\beta$ A1	302bp	For GCAGGAGCAGATGAGGAAAAGGGAG (SEQ ID NO:20) Rev CGCATGCGGTAGTGGTTGAT (SEQ ID NO:21)	1017...1094 1372...1253
INH $\beta$ A2	268bp	For GGCACGTCCGGTCCTCACTG (SEQ ID NO:22) Rev TCTTCATTTTGCCACTGTCTTCTC (SEQ ID NO:23)	1314...1334 1581...1558
<b>Inhibin Beta B</b>			
INH $\beta$ B	586bp	For CGTGGTGCCGGTGTTCTGTGGAC (SEQ ID NO:24) Ref CTCCACAGCCCAACAGAATGACT (SEQ ID NO:25)	617...638 1185...1163
INH $\beta$ B1	202bp	For CGTGGTGCCGGTGTTCTGTGGAC (SEQ ID NO:26) Rev GCCGGTGGGTGCTATGAT (SEQ ID NO:27)	617...638 1185...1163
INH $\beta$ B2	218bp	For GCACCCACCGGCTACTACG (SEQ ID NO:28) Rev TCCCGCTTGACGATGTTGT (SEQ ID NO:29)	807...825 1024...1006
INH $\beta$ B3	241bp	For AACTCCTGCTGCATTCCACCAA (SEQ ID NO:30) Rev CTCCACAGCCCAACAGAATGACT (SEQ ID NO:31)	945...967 1185...1163